The MRP4/ABCC4 Gene Encodes a Novel Apical Organic Anion Transporter in Human Kidney Proximal Tubules: Putative Efflux Pump for Urinary cAMP and cGMP

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Abstract. The cyclic nucleotides cAMP and cGMP play key roles in cellular signaling and the extracellular regulation of fluid balance. In the kidney, cAMP is excreted across the apical proximal tubular membrane into urine, where it reduces phosphate reabsorption through a dipyridamole-sensitive mechanism that is not fully understood. It has long been known that this cAMP efflux pathway is dependent on ATP and is inhibited by probenecid. However, its identity and whether cGMP shares the same transporter have not been established. Here the expression, localization, and functional properties of human multidrug resistance protein 4 (MRP4) are reported. MRP4 is localized to the proximal tubule apical membrane of human kidney, and membrane vesicles from Sf9 cells expressing human MRP4 exhibit ATP-dependent transport of [3H]cAMP and [3H]cGMP. Both probenecid and dipyridamole are potent MRP4 inhibitors. ATP-dependent [3H]methotrexate and [3H]estradiol-17β-d-glucuronide transport by MRP4 and interactions with the anionic conjugates S-(2,4-dinitrophenyl)-glutathione, N-acetyl-(2,4-dinitrophenyl)-cysteine, α-naphthyl-β-d-glucuronide, and p-nitrophenyl-β-d-glucuronide are also demonstrated. In kidneys of rats deficient in the apical anionic conjugate efflux pump Mrp2, Mrp4 expression is maintained at the same level. It is concluded that MRP4 is a novel apical organic anion transporter and the putative efflux pump for cAMP and cGMP in human kidney proximal tubules.

cAMP and cGMP are second messengers involved in the early response to extracellular signals in virtually every living cell. In many epithelial cells, they also play important roles in the extracellular regulation of fluid balance. It has been known for almost 30 yr that cAMP and cGMP are excreted from the kidney into urine (1). The main source of urinary cAMP seems to be kidney proximal tubules. Stimulation by parathyroid hormone results in elevation of intracellular cAMP levels and subsequent excretion of cAMP into the tubular lumen, where it is converted to adenosine (2). Reuptake of adenosine into proximal tubule cells reduces phosphate reabsorption via an unknown mechanism (3). The relevance of this extracellular cAMP-adenosine pathway has been indicated by the finding that dipyriramole, which inhibits adenosine uptake, resolves urinary phosphate leakage among patients with primary hyperparathyroidism or hypophosphatemia (3,4). Urinary cGMP, which originates from glomeruli under the control of atrial natriuretic peptide (ANP) or nitric oxide, inhibits reabsorption of sodium and chloride in the distal tubules (5,6). Proximal tubules seem to lack ANP receptors; however, apical proximal tubule membrane vesicles from ANP-infused rat kidneys exhibit decreased sodium uptake (7). Also, ANP inhibits sodium reabsorption in the proximal tubule-derived cell line LLC-PK1, which can be mimicked by exogenous cGMP (8). The identity of the proximal tubule urinary excretion route for cAMP and cGMP has not yet been resolved.

Monolayers of LLC-PK1 cells release both cAMP and cGMP preferentially across the apical membrane, via a probenecid-sensitive, temperature-sensitive, energy-dependent transport mechanism (8,9). Such a profile matches a member of the multidrug resistance protein (MRP/ABCC) subgroup of the ATP-binding cassette (ABC) protein superfamily (10,11). Mrp2 (ABCC2) functions as an ATP-dependent anionic conjugate efflux pump in the apical membrane of renal proximal tubules, small intestinal villi, and hepatocytes (12). The substrate specificity of MRP2, including leukotriene C4 (LTC4), S-(2,4-dinitrophenyl)-glutathione (DNP-SG), and estradiol-17β-d-glucuronide (E217βG), exhibits considerable overlap with that of Mrp1, which does not interact with cAMP (12,13). Furthermore, rats deficient in functional Mrp2 (Eisai hyperbilirubinemic Spague-Dawley [EHBR]/transport-deficient WistarTR− rats) do not display a clear defect in the renal excretion of several organic anions, suggesting the presence of a compensating apical organic anion transporter in the kidney (14). MRP5 (ABCC5) has been identified as a probenecid-
sensitive, ATP-dependent transporter for cGMP and cAMP but is localized to the basolateral membrane upon expression in MDCKII cells (15,16). MRP4 (ABCC4) exhibits the closest resemblance to MRP5, on the basis of structural characteristics, and both proteins mediate resistance to several xenobiotic nucleosides, such as the antiviral therapeutic agent 9-(2-phosphonomethoxyethyl)adenine (adefovir) (16–18). Pharmacokinetic studies with human subjects have indicated that the kidney is the main excretory pathway for adefovir and cidofovir [(5)-1-[(3-hydroxy-2-phosphonomethoxypropyl)]cytosine] (19). Furthermore, the excretory path is probencid-sensitive and rate-limiting, which explains the nephrotic side effects observed with these drugs (19).

In this report, we provide evidence that MRP4 is a novel renal proximal tubule apical organic anion efflux pump for cAMP and cGMP. We demonstrate the expression and immunolocalization of MRP4 in human kidney and the substrate specificity of recombinant human MRP4 by functional expression.

Materials and Methods

Materials

3H-labeled methotrexate (MTX) (15 Ci/mmol); [8-3H]cGMP (6.8 Ci/mmol); [2,8-3H]cAMP (21.9 Ci/mmol), and [3H]adefovir [9-(2-phosphonomethoxyethyl)adenine (adefovir) (16–18). Pharmacokinetic studies with human subjects have indicated that the kidney is the main excretory pathway for adefovir and cidofovir [(5)-1-[(3-hydroxy-2-phosphonomethoxypropyl)]cytosine] (19). Furthermore, the excretory path is probencid-sensitive and rate-limiting, which explains the nephrotic side effects observed with these drugs (19).

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Isolation of Full-Length Human MRP4 and Partial Rat MRP4 cDNA

Total RNA was isolated from human kidney cortex by acid guanidinium isothiocyanate-phenol-chloroform extraction and was reverse-transcribed in 50 mM Tris-HCl (pH 8.3), 75 mM KCl, 10 mM dithiothreitol, 3 mM MgCl2, supplemented with 0.5 mM dNTP, 10 μM reverse primer (see below), and 200 U of Moloney murine leukemia virus reverse transcriptase (Life Technologies, Breda, The Netherlands). For amplification of two overlapping fragments, primers were designed on the basis of the published human sequence (nucleotides 1 to 3979; GenBank accession number AF071202). The forward primer M1, KK 5'-GGTACGCCACACCTGCCCAGTG-3'; the Kpnl site is underlined, a partial Kozak sequence (21), introduced for future studies, is in bold, and ATG is italic) and the reverse primer M2, R2 5'-GGCTTCCAGACGCA-CATCTTTCAAGG-3', nucleotides 2030 to 4010) were used to amplify the 5' cDNA fragment. For amplification of the 3' cDNA fragment, the forward primer M1,F2 (5'-TGGTGCAGAAGGGGACTTACACTGAGTTC-3', nucleotides 1832 to 1860) and the reverse primer M1,R3 5'-CTTCGCCGACTGGAGGTAGTTT-3', nucleotides 3987 to 4010) were used. PCR was performed with the Advantage-2 PCR enzyme system (Clontech, Leusden, The Netherlands), which includes a nuclease-deficient Taq DNA polymerase and a proofreading polymerase. Amplifications were performed for 30 cycles of 94°C for 45 s, 55°C for 30 s, and 72°C for 3 min. The PCR fragments obtained were purified and cloned into pGEM-T Easy (Promega, Leiden, The Netherlands). Multiple clones were analyzed by cycle sequencing using an ABI 377 automated sequence (Applied Biosystems, Foster City, CA). The coding sequence was identical to the published sequence (GenBank/EMBL accession number AF071202) except for two 1-bp alterations (guanine for adenine at nucleotide positions 2030 and 3800), which resulted in substitution of arginine for glutamine at position 677 and glycine for serine at position 1267. A full-length human MRP4 cDNA in the EcoRI and SpeI sites of pGEM-T-Easy (pGTE-MRP4) was obtained by assembling the 5' (nucleotides 1 to 993) and 3' (nucleotides 1993 to 4010) cDNA fragments.

Total RNA was isolated from rat kidney cortex and reverse-transcribed with 10 μM oligo(dT) as described above. PCR with the human MRP4 primers described by Kool et al. (22) was performed for 35 cycles of 94°C for 30 s, 30°C for 1 min, and 72°C for 1 min. The resulting 239-bp fragment was cloned into the Smal site of pBlue-script KS (pBS-Mrp4) and sequenced as described above. The partial rat MRP4 cDNA has been submitted to GenBank/EMBL (accession number AF376781).

Antibodies

The linker region (amino acids 611 to 676) of human MRP4 was chosen as the epitope for a polyclonal antibody (pAb). With the M1,F2 and M2,R2 primers, the encoding cDNA was amplified from pGTE-MRP4 and cloned in-frame into the expression vector pGEX-3x (Pharmacia, Woerden, The Netherlands). A specific rat Mrp4 cDNA, which encodes a protein with 93% identity to amino acids 1165 to 1231 of human MRP4, was amplified from pBS-Mrp4 with the forward primer 5'-TAAATAATGGACACTGAGTTC-3' and the reverse primer 5'-AATGGTGAGAACAGTGCA-3'. The resulting 239-bp fragment was cloned into pGEX-2T. Glutathione S-transferase (GST) fusion proteins were expressed in DH5α Escherichia coli cells and isolated by glutathione-Sepharose 4B (Amersham, Uppsala, Sweden) affinity chromatography. Rabbits were immunized and boosted three times with fusion protein (400 and 200 μg, respectively), followed by collection of serum, which was affinity-purified with a column of immobilized GST and then a column of immobilized GST-fusion protein. The eluted, affinity-purified, anti-rat Mrp4 (pAb rM4-p1; 2 mg/ml) and anti-human Mrp4 (pAb hM4-p3; 3 mg/ml) antibodies were used in immunohistochemical and immunoblot analyses (see below). Monoclonal antibody (mAb) Mrp1,III-6 is directed against rat Mrp2 and cross-reacts with human MRP2 (23), and pAb G34 is directed against the rat sodium/phosphate type IIa cotransporter was kindly provided by Dr. J. Biber (Institute of Physiology, University of Zurich, Zurich, Switzerland).

Immunohistochemical Analyses

Normal tissue from a nephrectomized human kidney was provided by the Department of Pathology; its use for medical study was approved by the Nijmegen University institutional review board. Kidney slices were fixed in 1% (vol/vol) periodate-lysine-paraformaldehyde (PLP) fixative for 2 h, washed with 20% (vol/vol) sucrose in phosphate-buffered saline (PBS), and subsequently frozen in liquid N2.
cells were infected with recombinant baculovirus encoding the Sf9 cells were infected as described (26). For control experiments, Sf9 nant baculovirus encoding human MRP4 was generated by using the pFASTBAC1 multiple cloning site, to create pFB1-MRP4. Recombi- the full-length MRP4 cDNA was cloned into the /H11001 unit of rat H/K-ATPase (HKβ).

Expression of Human MRP4 in Sf9 Cells

From the construct pGTE-MRP4, an EcoRI fragment containing the full-length MRP4 cDNA was cloned into the EcoRI site of the pFASTBAC1 multiple cloning site, to create pFB1-MRP4. Recombi- nant baculovirus encoding human MRP4 was generated by using the Bac-To-Bac baculovirus expression system (Life Technologies), and Sf9 cells were infected as described (26). For control experiments, Sf9 cells were infected with recombinant baculovirus encoding the β-subunit of rat H⁺/K⁺-ATPase (HKβ).

Preparation of Membrane Vesicles from Kidney and Sf9 Cells

Brush border membrane vesicles (BBMV) and basolateral membrane vesicles (BLMV) from rat and human kidney cortex were isolated as described (27). Membrane vesicles from Sf9-MRP4 and control cells were isolated as described (28).

Deglycosylation Study and Immunoblot Analyses

Membrane vesicles were treated with PNGase F according to the instructions provided by the manufacturer. Immunoblotting was performed as described previously (26). Blots were incubated with pAb rM4-p1 (2 μg/ml), pAb hM4-p3 (6 μg/ml), G34 antiserum (diluted 1:1,000), or M3 III-6 (diluted 1:10) according to the same protocol. Densitometric analyses were performed by using Image Pro Plus 3.0 (Media Cybernetics, Silver Spring, MD).

Transport Studies in Membrane Vesicles

Freshly prepared membrane vesicles (150 μg protein-equivalents) from Sf9 cells expressing MRP4 or HKβ were prewarmed at 37°C for 1 min and added to prewarmed transport buffer [10 mM MgCl₂, 40 mM 3-(N-morpholino)propanesulfonic acid-Tris, pH 7.0, 50 mM KCl] supplemented with an ATP-regenerating mixture (4 mM ATP, 10 mM creatine phosphate, and 100 μM creatine kinase) and a [³H]-labeled compound (28). At indicated times, samples were withdrawn from the reaction mixture, diluted in cold transport buffer, and filtered through NC45 filters. Samples from [³H]ladeovir uptake studies were filtered through ME25 filters. After being washed with 5 ml of transport buffer, filters were dissolved in liquid scintillation fluid for determination of bound radioactivity. In control experiments, ATP was replaced by the nonhydrolyzable analog β,γ-methylene-ATP (AMP-PCP). Net ATP-dependent transport was calculated by subtracting values measured in the presence of AMP-PCP from those measured in the presence of ATP.

Results

The localization of MRP4 in kidneys was investigated with single- and double-labeling immunofluorescence microscopic analyses of PLP-fixed human kidney sections. To this end, pAb were raised against the linker region of human MRP4 (pAb hM4-p3) and a region near the carboxy terminus of rat Mrp4 (pAb rM4-p1). Incubation of sections with affinity-purified pAb hM4-p3 demonstrated specific immunofluorescence labeling at the brush border (apical) membrane of proximal tubules (Figure 1, A to D). Identical results were obtained with affinity-purified pAb rM4-p1 on PLP-fixed rat kidney sections (data not shown). The integrity of the proximal tubules with MRP4-specific labeling was confirmed by phase-contrast micro-scopy (Figure 1, B and D). We observed specific staining of the glomeruli, but the signal intensity varied in different label- ing experiments and was lower than that observed for proximal tubule brush border membranes. Distal tubules located in the kidney cortex, which can be recognized on the basis of apical membranes with a characteristic absence of a microvillar structure, were devoid of staining (Figure 1, C and D). When kidney sections were incubated with the secondary antibody alone (Figure 1E), with preimmune serum, or with fusion protein-pretreated hM4-p3 antibodies (data not shown), no staining was observed. Double-labeling of human kidney sections with pAb hM4-p3 and mAb M3 III-6 demonstrated colocalization of MRP4 and MRP2 (Figure 1, F to H).

The expression of human MRP4 and rat Mrp4 in kidneys was further examined by immunoblotting of BBMV and BLMV from human and rat kidneys. Human MRP4 was detected as a single band of 170 kD in the BBMV fraction, using pAb hM4-p3 (Figure 2A). Identical results were obtained for rat Mrp4 by using pAb rM4-p1. To confirm the isolation procedure for membrane vesicle fractions, the expression of MRP2 and the Na⁺/K⁺-ATPase was investigated. As expected, human MRP2 (190 kD) was detected in the BBMV fraction, whereas the β-subunit of human Na⁺/K⁺-ATPase (55 kD) was detected in the BLMV fraction. Identical results were obtained with BBMV and BLMV from rat kidney (data not shown). Both pAb hM4-p3 and rM4-p1 detected recombinant MRP4 expressed in Sf9 cells (Figure 2B), but the size was smaller than that of MRP4 detected in human kidneys. This difference in size can be attributed to the lack of complex glycosylation in Sf9 cells, which was confirmed by treatment of BBMV and membrane vesicles from Sf9-MRP4 cells with PNGase F (Figure 2C). The molecular weight of MRP4 detected in Sf9-MRP4 cells and PNGase F-treated BBMV matches the size calculated for MRP4 cDNA.

To investigate whether MRP4 is a candidate transporter for cAMP and cGMP, uptake studies were performed with membrane vesicles isolated from Sf9-MRP4 and Sf9-HKβ cells. The uptake of 100 μM [³H]cAMP (Figure 3A) or 1 μM [³H]cGMP (Figure 3B) into membrane vesicles from Sf9-MRP4 cells was stimulated in the presence of ATP and was linear up to 12 min of incubation. The initial net MRP4-mediated, ATP-dependent uptake rates for [³H]cAMP (Figure 3C) and [³H]cGMP (Figure 3D) were 8 pmol/mg per min (eightfold higher than control values) and 78 fmol/mg per min (12-fold higher than control values), respectively. MRP4-me- diated, ATP-dependent [³H]cGMP transport was inhibited by probenecid (IC₅₀ < 100 μM) and cAMP (IC₅₀ < 250 μM)
Dipyridamole proved to be a strong inhibitor of MRP4 (Table 1). Adefovir and cidofovir are excreted into urine via a probenecid-sensitive transport mechanism (19), and MRP4 was recently demonstrated to mediate resistance to adefovir (17,18). However, MRP4-mediated, ATP-dependent [3H]cGMP trans-

Figure 1. Immunofluorescence microscopic analyses of multidrug resistance protein 4 (MRP4) in human kidney. Proximal tubules (*) surrounding the glomerulus (g) exhibited staining at the brush border membrane with polyclonal antibody (pAb) hM4-p3, directed against human MRP4 (A and C). Distal tubules (arrows) were negative for MRP4 expression. Phase-contrast microscopic analyses of sections identical to A and C are also shown (B and D, respectively). No staining was observed when the secondary antibody was used alone (E). Double-labeling of a human kidney section with pAb hM4-p3 (F) and monoclonal antibody M2-III-6 (G) was used for detection of MRP4 and MRP2, respectively. Superimposition demonstrates colocalization of MRP4 with MRP2 in the proximal tubule brush border membrane (H).
port was inhibited by neither adefovir nor cidofovir, even at concentrations 3 orders of magnitude higher than the substrate concentration (Table 1). Furthermore, uptake of [3H]adefovir into membrane vesicles from Sf9-MRP4 cells in the presence of ATP was not different from uptake in the presence of AMP-PCP (data not shown). GSH, glucuronate, or N-acetyl-cysteine was added as a factor to potentially stimulate MRP4 transport, as previously demonstrated for MRP2-mediated, ATP-dependent [3H]vinblastine transport (29). Neither of these conditions resulted in ATP-dependent uptake of [3H]adefovir by MRP4.

MTX is eliminated from the body largely via the kidneys, and it is a substrate for MRP2 (28). We investigated whether it is also transported by MRP4. In the presence of ATP, uptake of 0.1 μM [3H]MTX into membrane vesicles from Sf9-MRP4 cells was linear up to 15 min (Figure 4A). The net ATP-dependent [3H]MTX uptake rate was approximately 10-fold higher than that for Sf9-HKβ membrane vesicles. MRP4-mediated, ATP-dependent [3H]MTX uptake increased with increasing MTX concentrations. Fitting of the recorded values to the Michaelis-Menten equation yielded a Vmax value of 430 ± 110 pmol/mg per min and a Km value of 1.3 ± 0.2 mM (Figure 4B). Concentration-dependent inhibition of ATP-dependent [3H]MTX transport by MRP4 was observed for DNP-SG and NAc-DNP-Cys, both with IC50 values of approximately 10 μM (Table 2). MRP4 was not inhibited by LTC4 (Table 2), and Sf9-MRP4 membrane vesicles did not exhibit ATP-dependent uptake of [3H]LTC4 with concentrations of either 5 nM or 1 μM. MRP4 was also inhibited by the glucuronide conjugates α-naphthyl-β-D-glucuronide (IC50 < 100 μM), p-nitrophenyl-β-D-glucuronide (IC50 < 1000 μM), and E2-17βG (Table 2). Sf9-MRP4 membrane vesicles exhibited time- and ATP-dependent uptake of [3H]E2-17βG at a concentration of 500 μM (17.6 mCi/mmol), but uptake was slow (Figure 5). At a tracer concentration of 0.2 μM, we did not observe MRP4-mediated, ATP-dependent [3H]E2-17βG transport, although, under identical conditions, uptake was observed in membrane vesicles from Sf9 cells expressing human MRP2 (data not shown).

Comparative studies have indicated that the absence of Mrp2 in TR− rat kidneys has no significant implications for the renal excretion of several organic anions. For evaluation of the level of expression of Mrp4, the BBMV fractions of kidney cortex from three wild-type rats and three TR− rats were isolated and subjected to immunoblot analysis using pAb rM4-p1 (Figure 6). To exclude differences in the yields of BBMV from different rats, protein-equivalent values were normalized to the alkaline phosphatase activity. The level of expression of the apical sodium/phosphate type IIa cotransporter (NaPi-IIa) was included as an internal control. Densitometric analysis indicated that the Mrp4/NaPi-IIa cotransporter ratio in wild-type rats (0.64 ± 0.23) was not significantly different from that in TR− rats (0.74 ± 0.18).
Table 1. Effects of probenecid, dipyridamole, and nucleosides on MRP4-mediated, ATP-dependent, [3H]cGMP transport

<table>
<thead>
<tr>
<th>Compound</th>
<th>Concentration (µM)</th>
<th>[3H]cGMP Uptake (% of Control)</th>
</tr>
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<tr>
<td>No inhibitor</td>
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<tr>
<td>Probencid</td>
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<td></td>
<td>100</td>
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<td></td>
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<td>Dipyridamole</td>
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</tr>
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<td>Adefovir</td>
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<td></td>
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<td>97 ± 12</td>
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* Membrane vesicles from Sf9-MRP4 cells were incubated with 1 µM [3H]cGMP for 10 min at 37°C, in the absence (control) or presence of various compounds. Net ATP-dependent uptake was expressed as a percentage of control uptake. Data represent mean ± SEM values of three determinations. MRP4, multidrug resistance protein 4.

b P < 0.05.

c P < 0.01 (ANOVA with Bonferroni’s correction).

Table 2. Effects of glucuronide and glutathione conjugates on MRP4-mediated, ATP-dependent, [3H]MTX transport

<table>
<thead>
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<th>Compound</th>
<th>Concentration (µM)</th>
<th>[3H]MTX Uptake (% of Control)</th>
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</thead>
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<tr>
<td>No inhibitor</td>
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<td>DNP-SG</td>
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<td>25 ± 5 c</td>
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<tr>
<td>Estradiol-17β-D-glucuronide</td>
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* Membrane vesicles from Sf9-MRP4 cells were incubated with 0.1 µM [3H]MTX for 10 min at 37°C, in the absence (control) or presence of various compounds. Net ATP-dependent uptake was expressed as a percentage of control uptake. Data represent mean ± SEM values of three to six determinations. MTX, methotrexate; DNP-SG, S-(2,4-dinitrophenyl)-glutathione; NAc-DNP-Cys, N-acetyl (2,4-dinitrophenyl)-cysteine.

b P < 0.05.
c P < 0.01 (ANOVA with Bonferroni’s correction).

Figure 4. ATP- and concentration-dependent transport of [3H]methotrexate ([3H]MTX) by MRP4. (A) Time-dependent uptake of [3H]MTX (0.1 µM) into membrane vesicles from Sf9-MRP4 (squares) and Sf9-HKβ (triangles) cells at 37°C. Solid and open symbols represent uptake in the presence of ATP and AMP-PCP, respectively. (B) Incubation of Sf9-MRP4 membrane vesicles for 10 min at 37°C in the presence of [3H]MTX concentrations ranging from 0.1 µM to 5 mM. Net ATP-dependent uptake values were fitted according to the Michaelis-Menten equation. Data represent the mean ± SEM of three determinations.

Discussion

The release of cAMP and cGMP in urine is an important step in their functioning as paracrine modulators of renal fluid homeostasis. It has been suggested that these endogenous nucleotides reach the tubular fluid via an ATP-dependent organic anion transporter (3). However, the identity of this transporter has not been established, and it is not known whether cAMP and cGMP share the same export pathway. In this report, we clearly demonstrate, using immunohistochemical and immunoblot analyses, that MRP4 is expressed at the apical (brush border) membrane of human kidney proximal tubules and that MRP4 mediates ATP-dependent transport of [3H]cAMP and [3H]cGMP. Furthermore, we demonstrate the interaction of MRP4 with various anionic conjugates and MTX.

The localization of MRP4 and its ability to transport cAMP and cGMP are in agreement with the results of previous functional studies that indicated the release of cAMP from renal proximal tubule cells upon stimulation by parathyroid hormone, leading to intracellular cAMP levels of at least 100 µM (9). Urinary cAMP has been suggested to originate from plasma cAMP by uptake via basolateral organic anion transporter 1 (OAT1) (1,30). However, plasma cAMP concentrations are in the nanomolar range (31), which may be too low for transport by OAT1; furthermore, infused [3H]cAMP is almost completely filtered by the glomeruli (1). Excreted cAMP has been demonstrated to be directly involved in the inhibition of phosphate reabsorption, after degradation to adenosine and subsequent uptake into the proximal tubules (3). Dipyridamole inhibits adenosine uptake and can be used to protect patients from renal phosphate leakage (4). Our results indicate that dipyridamole may have dual effects, i.e., reduction of adenosine uptake and also inhibition of cAMP release via MRP4, reducing urinary adenosine availability. The source of urinary cGMP has been localized primarily to the glomeruli (6). However, modulation...
of sodium reabsorption by cGMP in proximal tubules has also been reported, as well as the role of extracellular cGMP as a modulator of sodium transport in LLC-PK₁ cells (7,8). The presence of MRP4 in proximal tubules, combined with the observed substrate specificity, suggests that this part of the nephron also contributes to urinary cGMP. Whereas a high but physiologic concentration of [³H]cAMP (100 µM) was used to demonstrate transport by MRP4, a much lower concentration of [³H]cGMP (1 µM) was required. These results are in line with previous observations indicating that intracellular levels of cGMP are approximately 20-fold lower than cAMP levels and that efflux of cGMP from cells is significantly more efficient, compared with that of cAMP (31,32). High concentrations of cAMP were needed to inhibit ATP-dependent [³H]cGMP transport by MRP4 (this study), similar to findings for MRP5 (although with a lower IC₅₀ value) (15). During the final preparation of this manuscript, a study by Chen et al. (33), in which these cyclic nucleotides were identified as MRP4 substrates, was reported. Further studies with polarized monolayers of proximal tubule cells will be required to confirm that MRP4 is the renal proximal tubule efflux pump for cAMP and cGMP.

MRP4 and MRP5 have recently been associated with cellular resistance against adefovir (17,18). Furthermore, adefovir and cidofovir accumulate in proximal tubule cells, causing nephrotoxicity (19). The proximal tubule basolateral uptake carrier OAT1 was recently demonstrated to transport both adefovir and cidofovir (34). Pharmacologic studies with human subjects have suggested that the apical efflux pump, rather than the uptake carrier, represents the probenecid-sensitive, rate-limiting step in the urinary excretion of these compounds (19). Those data prompted us to speculate that MRP4 is the renal efflux pump for adefovir and cidofovir. However, those compounds did not inhibit MRP4-mediated transport, and we did not observe any evidence of ATP-dependent uptake of [³H]adefovir in membrane vesicles from MRP4-expressing Sf9 cells. It therefore remains unclear how MRP4 extrudes xenobiotic monophosphate nucleosides from the cell. Adefovir and cidofovir are monophosphates and are metabolized into di- and triphosphate (the active form) derivatives (35). HPLC analysis has suggested that MRP4 transports only the unmetabolized derivative and that both adefovir and cidofovir are excreted unchanged into the urine (17,19). However, the possibility that the di- or triphosphate metabolite is the actual substrate, which may be difficult to detect because of rapid degradation by extracellular enzymes, cannot be excluded. In the case of cidofovir, a cyclic metabolite has been identified in urine (36). With MRP4 being an efflux transporter for endogenous cyclic nucleosides, the cyclic metabolite, rather than cidofovir itself, may be transported. However, MRP4 and MRP5 may require a cofactor that is as yet unknown. In a similar manner, both MRP1 and MRP2 require GSH for ATP-dependent transport of the anticancer agents vincristine, vinblastine, and daunorubicin (29,37). Our results at least do not support a role for GSH in stimulating drug transport by MRP4.

Rats deficient in functional Mrp2 do not display clear differences in the renal excretion of many organic anions, compared with wild-type rats (14). Furthermore, transport studies with killifish renal proximal tubules have provided evidence for organic anion transporters in addition to MRP2 (38,39). We demonstrate that the substrate specificity of MRP4 overlaps...
with that of MRP2. The MRP2 substrates E17βG and MTX are both transported by MRP4, and the affinity of MRP4 for MTX (Km of approximately 1 mM) is in the same range as that documented for MRP2 (28). However, E17βG, a high-affinity MRP2 substrate (Km of approximately 7 µM), seems to be a low-affinity substrate for MRP4, because a concentration of 500 µM was required for detection of [3H]E17βG transport by MRP4 and 100 µM E17βG inhibited MRP4-mediated [3H]MTX transport by only 25%. On the basis of the results of inhibitor studies, we propose that α-naphthyl-β-D-glucuronide, p-nitrophosphoryl-β-D-glucoside, DNP-SG, and NAC-DNP-Cys are substrates for MRP4. Of interest, the excretion of α-naphthyl-β-D-glucuronide from TR− rabbit kidneys was found not to be different from that from wild-type kidneys (40). In rat kidneys, three additional organic anion transporters, i.e., organic anion transporter polypeptide 1 (Oatp1), kidney-specific organic anion-transporter 1 (Oat-k1), and sodium/phosphate cotransporter type 2 (Npt1), are expressed at the proximal tubule brush border membrane, with substrate specificities overlapping those of MRP2 and MRP4 (14,41). A human ortholog has been identified for rat Npt1; however, human counterparts of rat Oatp1 and Oat-k1 do not seem to exist (41,42). Whether rat MRP4 in TR− rats and human MRP4 in patients with Dubin-Johnson syndrome (both lacking functional MRP2) (12) can represent an alternative pathway for the renal excretion of organic anions remains to be established and may even involve differences between species.

References


